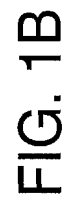


FIG. 1

FIG. 1A



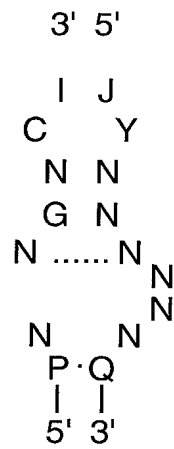


FIG. 2

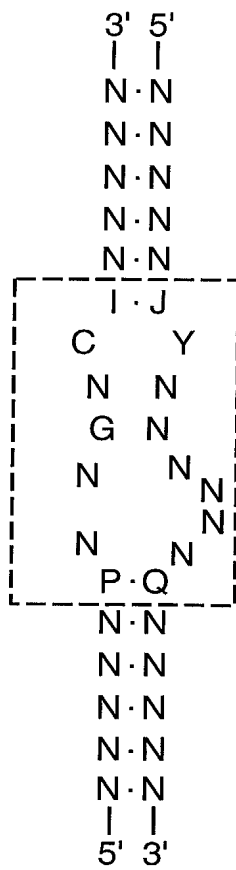
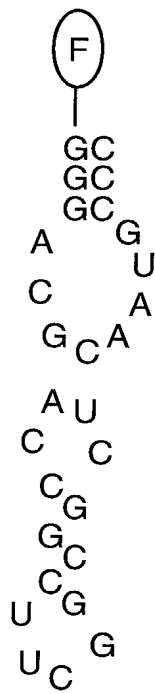


FIG. 4



HCV mIRES RNA

FIG. 6A

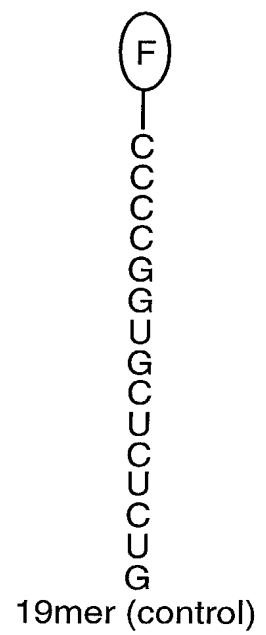


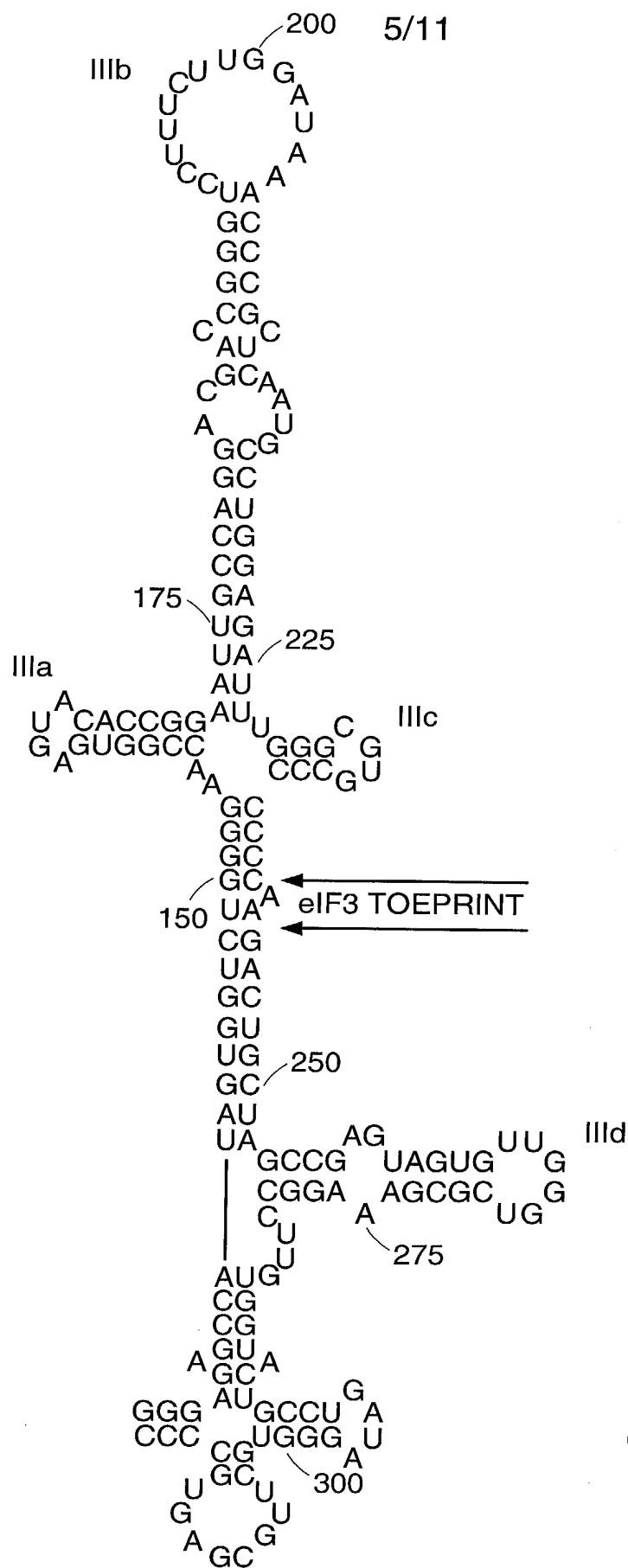
FIG. 6B

gi	1009264	GACGACC
gi	1160327	-----
gi	1438429	-----
gi	1181831	-----
gi	1197128	--A---T
gi	1805843	-GT----
gi	1930117	-GT----
gi	1066559	--A---T
gi	1066561	--T----
gi	1805796	-----
gi	329915	--T----
gi	1066557	-----
gi	1066567	-----
gi	2388526	--T----
gi	1066617	--T----
gi	1930119	--T----
gi	1183032	-TT----
gi	1197096	--A---T
gi	2809127	-----
gi	1009258	--T----
gi	1009266	-----
gi	1066565	-----
gi	1066577	-----
gi	1805846	-----
gi	1805851	-----
gi	1805853	A-A---T
gi	2465799	-GT----
gi	2605601	-----
gi	2809126	-----
gi	2809128	-----
gi	296246	--A---T
gi	3152988	-CT----
gi	3152993	--T----
gi	3152997	-----
gi	474334	--T----

FIG. 3A

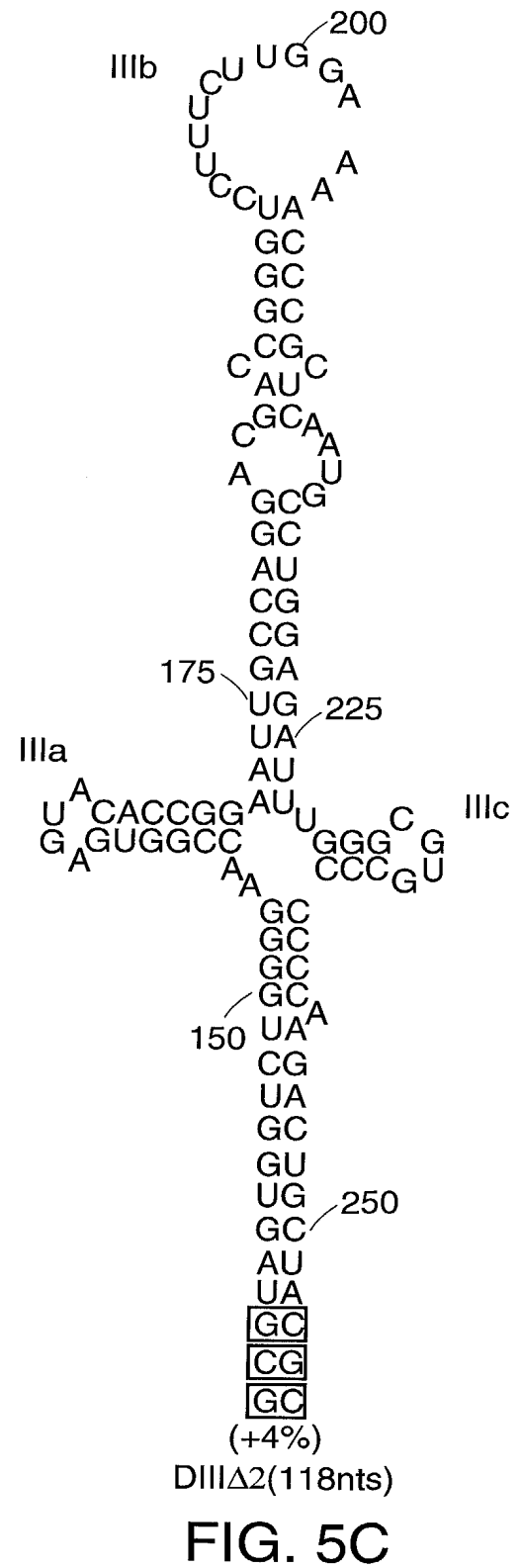
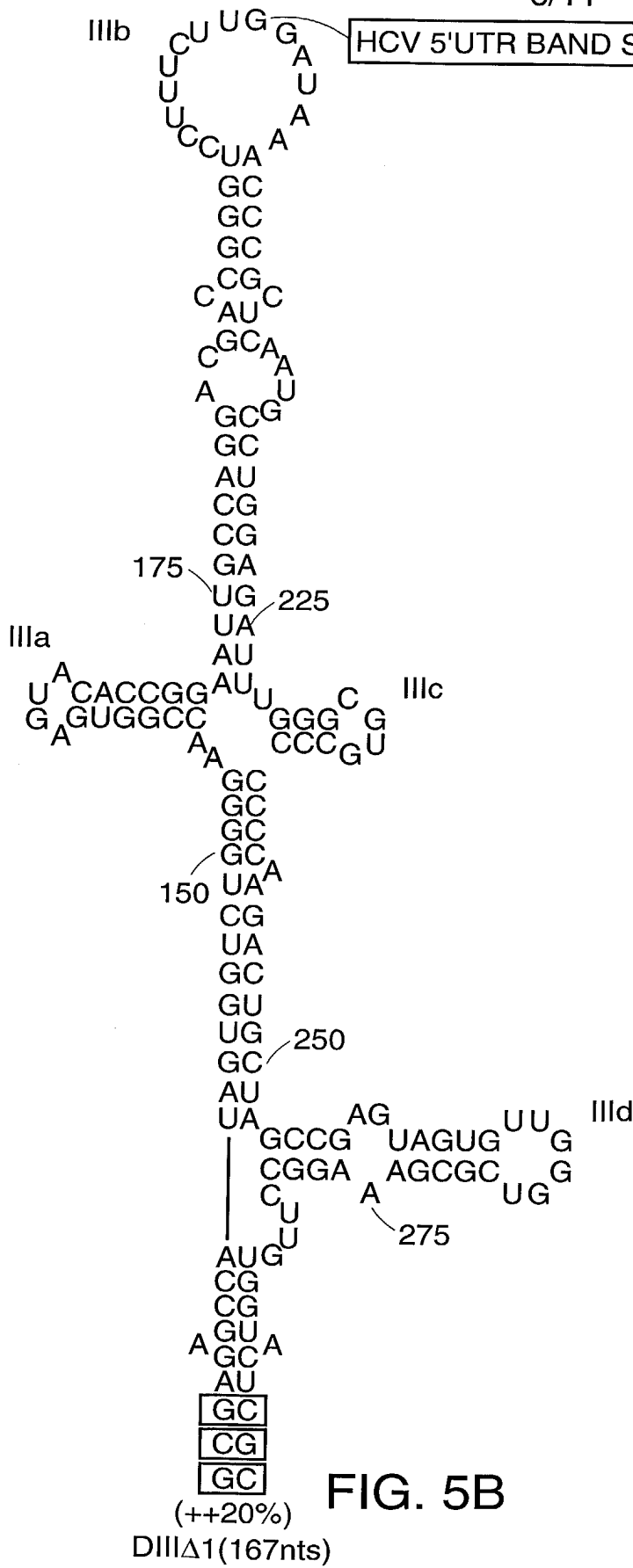
gi	1009264	GCTCAA	TGC
gi	1160327	-----	---
gi	1438429	-----	---
gi	1181831	-----	---
gi	1197128	A---T-	---
gi	1805843	-----	-A-
gi	1930117	-----	-A-
gi	1066559	A---T-	---
gi	1066561	-----	---
gi	1805796	-----	---
gi	329915	-----	---
gi	1066557	-----	---
gi	1066567	-----	---
gi	2388526	-----	---
gi	1066617	-----	---
gi	1930119	-----	---
gi	1183032	-----	---
gi	1197096	A---T-	---
gi	2809127	-----	---
gi	1009258	-----G	---
gi	1009266	-----	---
gi	1066565	-----	---
gi	1066577	-----	---
gi	1805846	A-----	---
gi	1805851	----G-	---
gi	1805853	A---T-	--T
gi	2465799	-----	-A-
gi	2605601	-----	---
gi	2809126	-----	---
gi	2809128	-----	---
gi	296246	A---TG	---
gi	3152988	-----	-A-
gi	3152993	-----	---
gi	3152997	-----	---
gi	474334	-----	---

FIG. 3B



eIF3(+++100%)
DIII(187nts)
(W/T+MUTANTS)

FIG. 5A



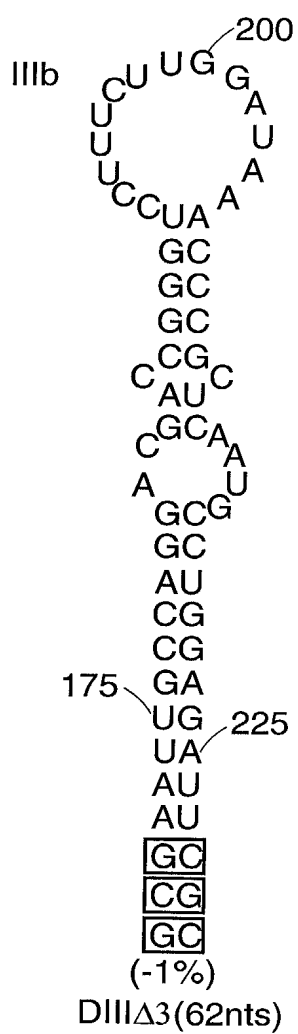


FIG. 5D

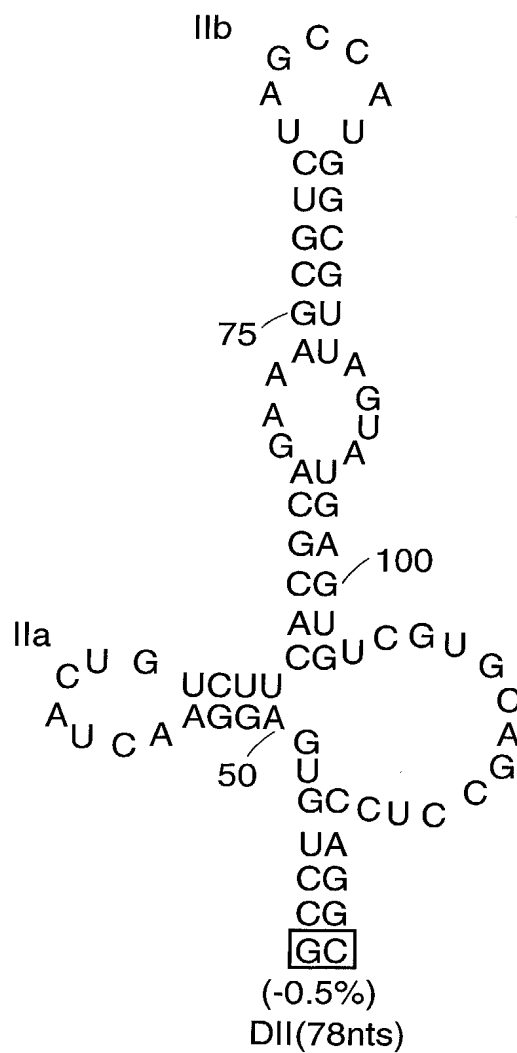


FIG. 5E

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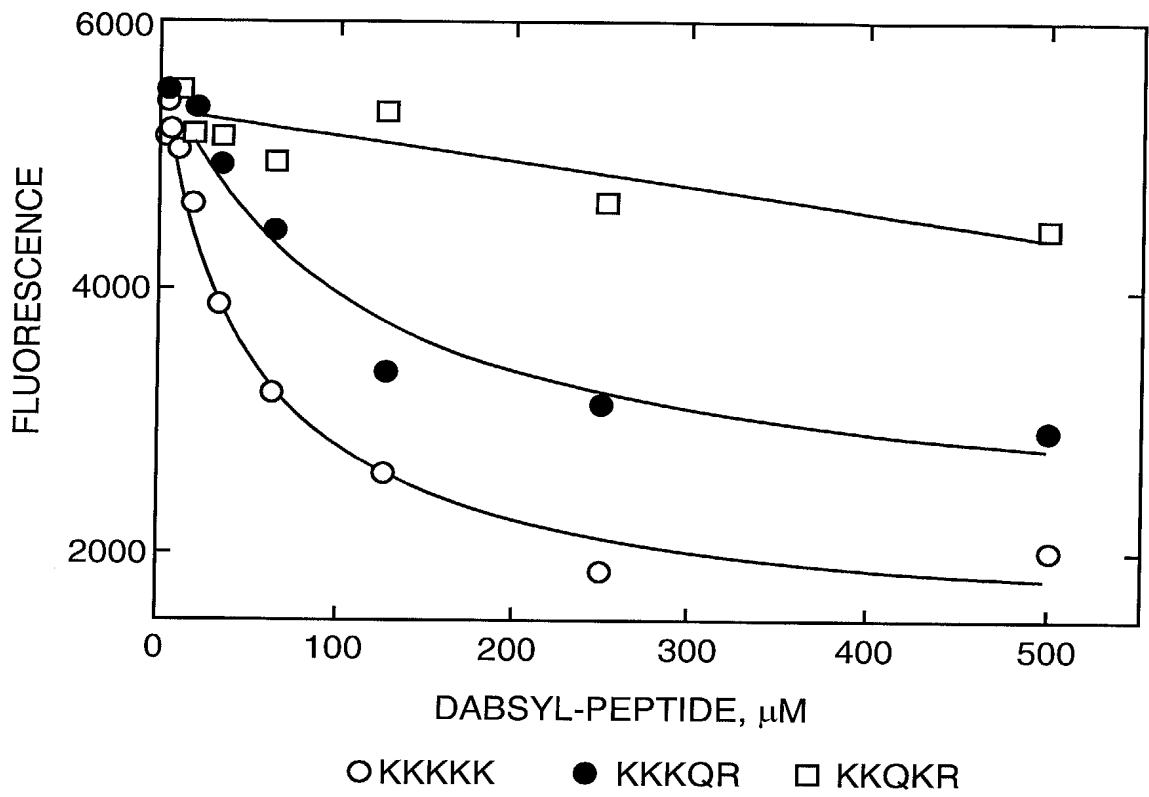


FIG. 7A

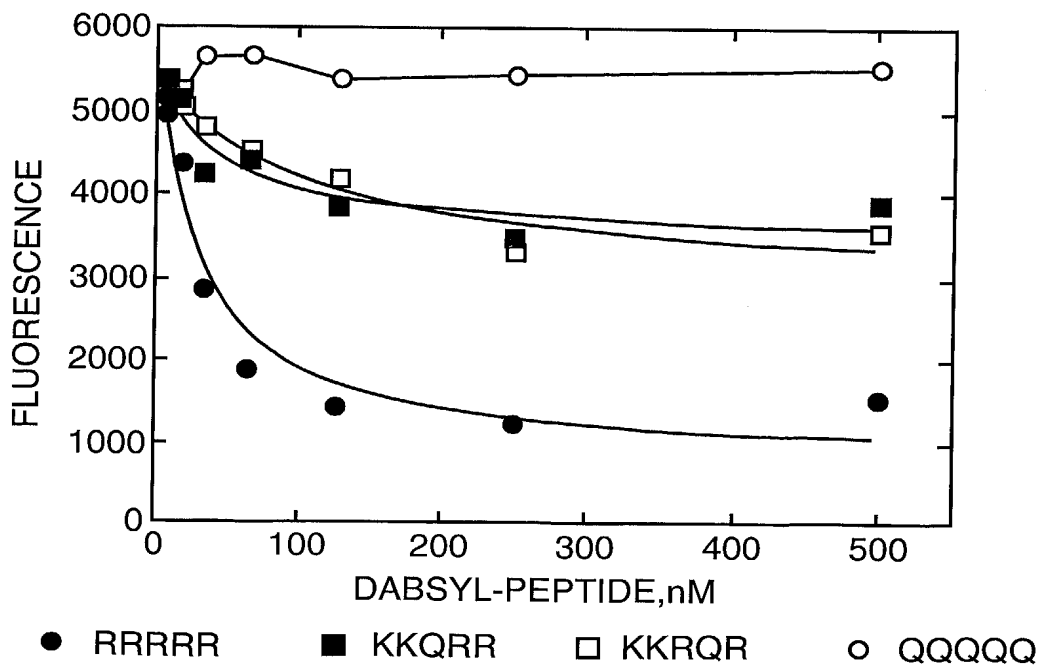


FIG. 7B

PAROMOMYCIN-TAMRA BINDING TO HCV IIIB

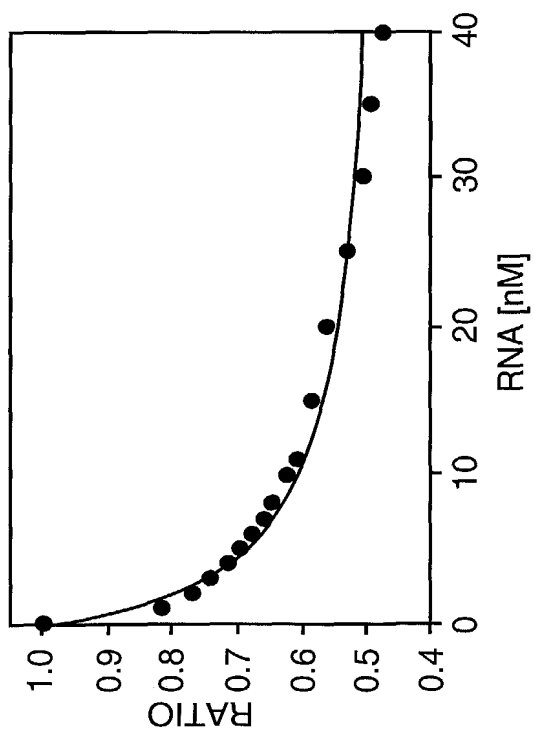


FIG. 8A

COMPETITION BY NEOMYCIN

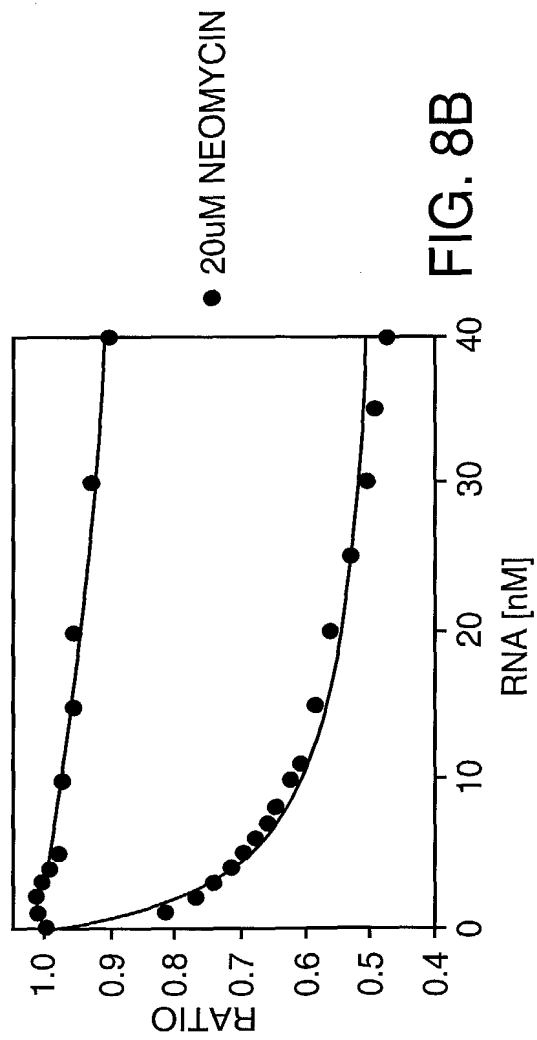


FIG. 8B

Diagram illustrating the secondary structure of the 16S rRNA gene of *E. coli*. The structure is shown as a large loop with a dashed box highlighting a specific region. The sequence within the dashed box is:

```

    C U A A
    GC
    GC
    GC
    C C C
    C A U C
    C G C A
    A U
    G C G
    G C
    A U
    C G
    C G
    G A
  
```

U A CACCGG AU AU U C IIIc
G GUGGCC GGG G U
A A CCC G
GC

GC
GC
CG
GC^A
UA
CG
UA
GC
GU
UG
GC²⁵⁰
AU
UA

275

2

①

FIG. 8A-1B

FIG. 8A-1A

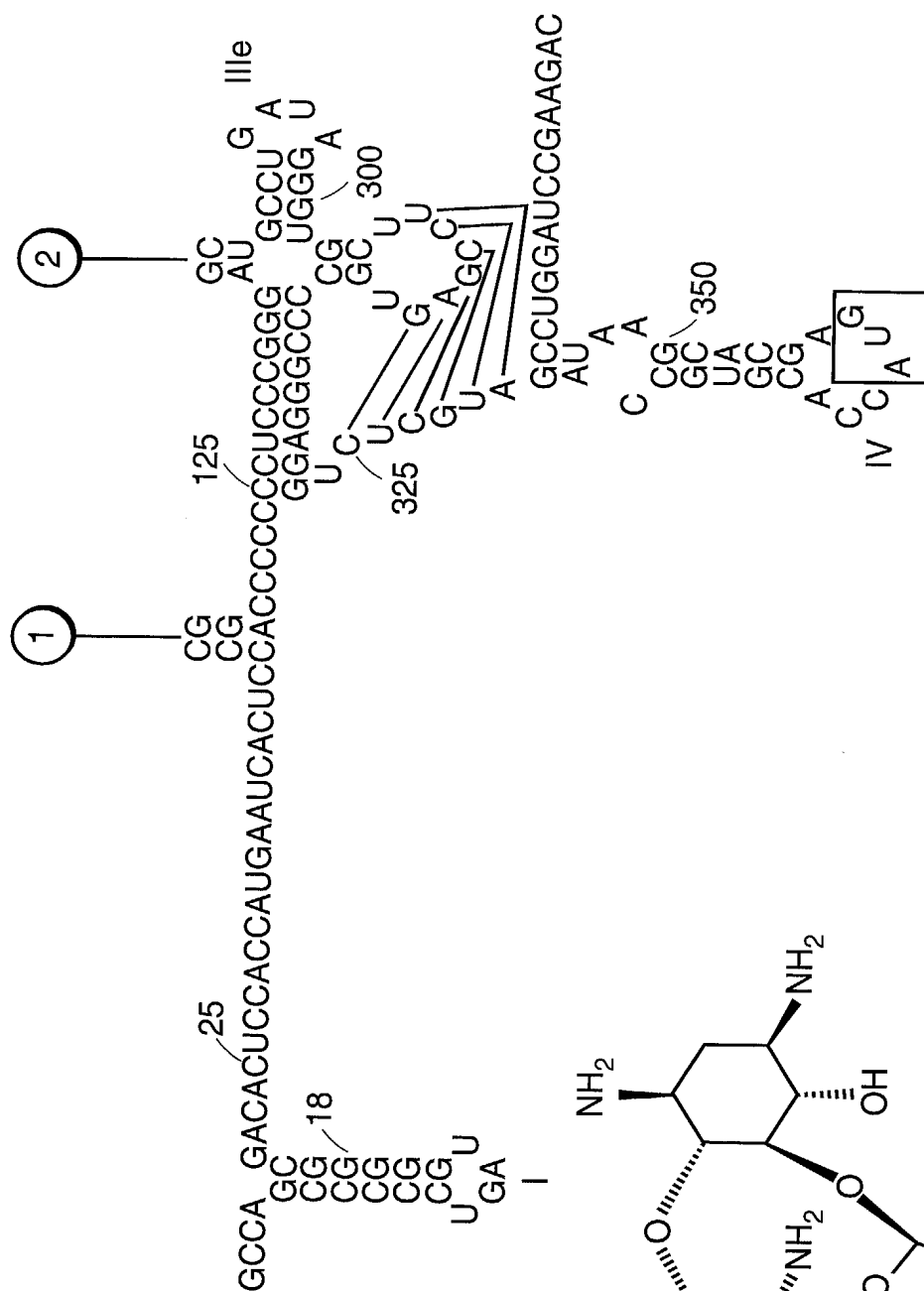
[illegible]

FIG. 8A-1B

